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COHEN, Daniel  
ESSIOUX, Laurent

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00416384-101290

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**09-06-2017**

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Gln	Phe	Asn	Cys	Arg	Lys	Thr	Ile	Pro	Cys	Lys	Gln	Tyr	Cys	Leu	Glu
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Tyr	Gly	Gly	Leu	Ser	Ser	Phe	Ile	Cys	Thr	Gly	Leu	Tyr	Glu	Thr	Phe
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Gly Ala Trp Met Cys Arg Gln Tyr Asp Asp Gly Leu Lys Ile Trp Leu
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Ala Ala Pro Arg Glu Asn Glu Lys Pro Phe Ile Asp Ser Glu Arg Ala
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cag aaa tgg cga ctg tct ctg gct tct ctc ttg ttt ttc aca gtc ctg      202
Gln Lys Trp Arg Leu Ser Leu Ala Ser Leu Leu Phe Phe Thr Val Leu
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Arg Asp Lys Glu His His Gln Gln Gln Gln Gln Gln Gln Gln Gln
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Arg Leu Leu Ser Ala Ser Ser Ser Pro Thr Leu Pro Pro Ser Pro Gly
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Gly Gly Gly Gly Ser Lys Gly Asn Arg Gly Lys Asn Asn Arg Ser Arg
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Cys Tyr Pro Gln Gly Ala Ser Ser Gly Gln Cys Phe Thr Val Glu Ser
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gag gag cag tcg tcc agg ggc tct cgg cca act ccg ctg tgg aac ttg      730
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09416384-101299

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Phe	Ser	Gly	Leu	Ser	Ser	Pro	Ser	Thr	Leu	Asn	Cys	Ser	Leu	Asp	Val	
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gtg	ctc	acg	gag	ggc	ggg	gag	atg	acc	acg	tgt	aga	cag	tgc	atc	gag	874
Val	Leu	Thr	Glu	Gly	Gly	Glu	Met	Thr	Thr	Cys	Arg	Gln	Cys	Ile	Glu	
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Ala	Tyr	Gln	Asp	Tyr	Asp	His	His	Ala	Gln	Glu	Lys	Tyr	Glu	Glu	Phe	
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Glu	Ser	Val	Leu	His	Lys	Tyr	Leu	Gln	Ser	Asp	Glu	Tyr	Ser	Val	Lys	
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Ser	Cys	Pro	Glu	Asp	Cys	Lys	Ile	Val	Tyr	Lys	Ala	Trp	Leu	Cys	Ser	
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Lys	Gln	Tyr	Cys	Leu	Glu	Val	Gln	Thr	Arg	Cys	Pro	Phe	Ile	Leu	Pro	
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Gly	Leu	Tyr	Glu	Thr	Phe	Leu	Thr	Asn	Asp	Glu	Pro	Glu	Cys	Cys	Asp	
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Ile	Arg	Ser	Glu	Glu	Gln	Thr	Ala	Pro	Arg	Pro	Lys	Gly	Thr	Val	Asp	
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Arg	Arg	Asp	Ser	Cys	Pro	Arg	Thr	Ser	Leu	Thr	Val	Ser	Ser	Ala	Thr	
		405				410					415					
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420					425					430					435	
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Leu	His	Thr	Val	Leu	Thr	Ala	Ser	Ala	Ala	Gln	Asn	Ser	Thr	Gly	Leu	
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Gly	Leu	Gly	Gly	Leu	Pro	Thr	Leu	Glu	Asp	Asn	Ser	Thr	Arg	Glu	Asp	
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Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Gln	Arg
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Gln	Ala	His	Arg	Leu	Leu	Ser	Ala	Ser	Ser	Ser	Pro	Thr	Leu	Pro	Pro
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Arg	Ser	Arg	Ala	Leu	Phe	Leu	Gly	Asn	Ser	Ala	Lys	Pro	Val	Trp	Arg
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Leu	Glu	Thr	Cys	Tyr	Pro	Gln	Gly	Ala	Ser	Ser	Gly	Gln	Cys	Phe	Thr
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Val	Glu	Ser	Ala	Asp	Ala	Val	Cys	Ala	Arg	Asn	Trp	Ser	Arg	Gly	Ala
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Trp	Asn	Leu	Ser	Asp	Phe	Tyr	Leu	Ser	Phe	Cys	Asn	Ser	Tyr	Thr	Leu
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	370					375					380				
Cys	Cys	Asp	Ile	Arg	Ser	Glu	Glu	Gln	Thr	Ala	Pro	Arg	Pro	Lys	Gly
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662107-10125

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taaaacatgt	aaactctaag ctctctcctt atagtcagtt agtagcagca catcttaaaa 300
tctgggtgtga	atattctcct agttctacat gagtctaact aaacagagga ttattcttag 360
gtgtttgaaa	gagacatatg tgacactgct gttttgagaa caatttaagt gttgtcttgt 420
catgtacaga	agttctcata ttactttaca taaatggttg cataattggt ttatagtata 480
taatagactg	tcaatatttc taggataact ccaaaacaaa atttcctaga mmacattttg 540
aaaagg	

<210> 35  
 <211> 502

<212> DNA  
 <213> Homo sapiens  
 <220>  
 <221> allele  
 <222> 185  
 <223> 99-15664-185 : polymorphic base G or T

<220>  
 <221> misc\_binding  
 <222> 162..208  
 <223> 99-15664-185.probe

<220>  
 <221> primer\_bind  
 <222> 166..184  
 <223> 99-15664-185.mis

<220>  
 <221> primer\_bind  
 <222> 186..204  
 <223> 99-15664-185.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..19  
 <223> 99-15664.pu

<220>  
 <221> primer\_bind  
 <222> 483..502  
 <223> 99-15664.rp complement

<220>  
 <221> misc\_feature  
 <222> 54  
 <223> n=a, g, c or t

<400> 35  
 gtttaccatt agcactgtca tatttgtgtg acttgtcatt ctctacagcg gagnacgggc 60  
 tggcacgggg cctgatgctg acttgcacaa gggaagcctc ctgtctctga cttccccagg 120  
 ataattcctg gggaaagtgt gctccctagt gttaagagcg gtttaatggc tggaggggtt 180  
 cagckggctg accaggcaga gaaggagggt gaatcacctc tcagcactct ccacttagac 240  
 tttgtgtggt cgtcgggtgg tcaaaccctc taactagtgtg tattgcagat ttggcattcc 300  
 agtgcaaaca aaagacagaa acacaatggt cacatgcttt ccagagatca cctggatata 360  
 agatcatttg attttcaagt aagtcgaaac cttggtggaa atcattaact atcctgttta 420  
 tgacaaaaaa ataaaatccc aaatttcttc tcttcatttc ttacctgctt taaaattgta 480  
 tccaaagcgt graagtaaaa ga 502

<210> 36  
 <211> 455  
 <212> DNA  
 <213> Homo sapiens



<220>  
 <221> allele  
 <222> 205  
 <223> 99-5919-215 : polymorphic base A or G

<220>  
 <221> misc\_binding  
 <222> 182..228  
 <223> 99-5919-215.probe

<220>  
 <221> primer\_bind  
 <222> 186..204  
 <223> 99-5919-215.mis

<220>  
 <221> primer\_bind  
 <222> 206..224  
 <223> 99-5919-215.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..19  
 <223> 99-5919.pu

<220>  
 <221> primer\_bind  
 <222> 435..455  
 <223> 99-5919.rp complement

<400> 36  
 ctacagcaat gcagatttca attctgccat tgaattccca gacatattcg tcatcccat 60  
 tttcatcccc caccacctg ccattttctt cgtgttaact tgttttcctg actcacagaa 120  
 atcacctttt cctgtataca tttttaggat gtcagacttt attctaataga tttctcctag 180  
 ttgcccccca aaattgtatt ctacrgtgtg attttaaaagc tgaattttca agatgatatt 240  
 tcatatctat attttcacia gcttttcttc tatgaatgtt attgtcagct gtcagggtgt 300  
 gagatggtac ttgatactac attctttcca agctgttgcc tgaatcggtt taagacaaag 360  
 tcattactag gctgtaaact gttgctctgc aaaattgagc agcacgtatt taaccactca 420  
 tacttcttag ctctccaaca ctttgagtca ataga 455

<210> 37  
 <211> 450  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> allele  
 <222> 157  
 <223> 99-5862-167 : polymorphic base C or T

<220>  
 <221> misc\_binding  
 <222> 134..180  
 <223> 99-5862-167.probe

<220>  
 <221> primer\_bind  
 <222> 138..156  
 <223> 99-5862-167.mis

<220>  
 <221> primer\_bind  
 <222> 158..176  
 <223> 99-5862-167.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..20  
 <223> 99-5862.pu

<220>  
 <221> primer\_bind  
 <222> 430..450  
 <223> 99-5862.rp complement

<400> 37  
 aatcaaggta gagatgtatg agaaatagcc ggttaaagaa acagcattac tttcagacta 60  
 tottttattt gaaatacacg tggggaaacc agaagggtgaa accccttagg agatggatat 120  
 aggatactaa aatctgagtt agaaaaattt gagcatyagc accttacgtg tcatgctaag 180  
 atagtgaatg agactgcaca ggaattgcat gcagtttaac ggaaaaagaa gtcgaaagat 240  
 aaatttcctag aacactaaca ccgagttatg ggaggagaaa tatcctgcac aggtcactct 300  
 gggagacatg tcaattgttt agccaatatc catttaactc atctttcttc ctaatgaaaa 360  
 ccgaatttgg agaagcaggt agtgcccctg gctagaaata tgaaccttcc cagcttctct 420  
 catgcactga actgacaaag ttcaggtctg 450

<210> 38  
 <211> 403  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> allele  
 <222> 292  
 <223> 99-16032-292 : polymorphic base A or C

<220>  
 <221> misc\_binding  
 <222> 269..315  
 <223> 99-16032-292.probe

<220>  
 <221> primer\_bind  
 <222> 273..291  
 <223> 99-16032-292.mis

<220>  
 <221> primer\_bind  
 <222> 293..311

<223> 99-16032-292.mis complement

<220>

<221> primer\_bind

<222> 1..19

<223> 99-16032.pu

<220>

<221> primer\_bind

<222> 384..403

<223> 99-16032.rp complement

<400> 38

gttggttacc	cacttcttcc	ccccagctcc	cccttctctca	cacagttcat	gccacatgcc	60
actctcctgg	actactggaa	atgcgtcagt	ccactctggg	ctcatcccat	catcccccat	120
gctgcaacct	gagagagagt	tgcaagttgc	aaatctgata	ttgtcaccac	cactctccac	180
actaaatccc	tctaattgct	ccccctttct	ttttggataa	attccttctg	cttgcatagc	240
cacgtgggtg	gcttctatag	catcacttca	cactgtgggc	acctgccttc	tmctcactca	300
ggaacttctc	tccattgaag	aagttcttct	tccccatctc	cagggctttc	ccactgacag	360
ttgtatctcc	cccataccaa	gcccaggtgg	tcattctcatc	cca		403

<210> 39

<211> 476

<212> DNA

<213> Homo sapiens

<220>

<221> allele

<222> 118

<223> 99-16038-118 : polymorphic base A or G

<220>

<221> misc\_binding

<222> 95..141

<223> 99-16038-118.probe

<220>

<221> primer\_bind

<222> 99..117

<223> 99-16038-118.mis

<220>

<221> primer\_bind

<222> 119..137

<223> 99-16038-118.mis complement

<220>

<221> primer\_bind

<222> 1..19

<223> 99-16038.pu

<220>

<221> primer\_bind

<222> 456..476

<223> 99-16038.rp complement

<400> 39

gttgcttatt	ctttctctct	tctgcagggt	ataaaggaat	ctgaacacga	ctgatatttt	60
ctttaatttt	tagatccaga	tatacattgg	gtaaaatcta	cttcataagg	tttcaaarga	120
gcattcttct	gagcaaactc	gaaaactctc	taaactctat	tggtatgtta	ctctttatct	180
ttatatgaat	ttaaattctt	ctagaagtta	gataaaactg	tggtaaagct	acataatact	240
tttgacatat	tttcaagcgt	agacaaactt	caattaattt	gtaagataca	ggaagaaaat	300
ttttccagtt	aaaatgtacc	tcttgggttc	tgagtggtta	gcaaccattc	acacttacag	360
ttcaaacagt	gcaaccttgt	aaaacatata	taacttatga	agagatcgat	atctcttttt	420
ataaagcaaa	caagtaaatt	tttcctcaa	tccatgattt	atttttgtga	agtggg	476

<210> 40

<211> 498

<212> DNA

<213> Homo sapiens

<220>

<221> allele

<222> 133

<223> 99-5897-143 : polymorphic base A or C

<220>

<221> misc\_binding

<222> 110..156

<223> 99-5897-143.probe

<220>

<221> primer\_bind

<222> 114..132

<223> 99-5897-143.mis

<220>

<221> primer\_bind

<222> 134..152

<223> 99-5897-143.mis complement

<220>

<221> primer\_bind

<222> 1..18

<223> 99-5897.pu

<220>

<221> primer\_bind

<222> 475..492

<223> 99-5897.rp complement

<400> 40

aaaagtgttt	gccagtcctg	tttcttacag	agcacagaac	tcagatgctc	ttataaagat	60
acaggataaa	tcacatcatt	tcctgctcca	tcacacagaat	attattatat	gatttagatc	120
acttttttaa	aamagaacat	ggacttagta	cagaacaaca	gcaaaagcct	ggggaaggag	180
aggagtgcac	catgaggagt	caatggggag	cagaagccag	tccatttgac	tgatttggtt	240
cgtgtgcaaa	ataattgcta	aataattgca	tatatgtgag	actccgggta	ttttcaaaac	300
cagctggcaa	aattgtgtta	ttctctaccc	tctgctggct	ttcacgggtt	ctctgttctc	360



<220>  
 <221> allele  
 <222> 97  
 <223> 99-13925-97 : polymorphic base A or G

<220>  
 <221> misc\_binding  
 <222> 74..120  
 <223> 99-13925-97.probe

<220>  
 <221> primer\_bind  
 <222> 78..96  
 <223> 99-13925-97.mis

<220>  
 <221> primer\_bind  
 <222> 98..116  
 <223> 99-13925-97.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..20  
 <223> 99-13925.pu

<220>  
 <221> primer\_bind  
 <222> 513..533  
 <223> 99-13925.rp complement

<400> 42  
 catggaagta aaagcatatc ttcattataa gacttctaca caaattatca catctttact 60  
 tacagcagct gaaacctgga aacaactcta atgccrctca acagaggaat ggatggataa 120  
 agaaactgtg atgcagtggg atacgactca acgaagatga gactaaaaat aattatactg 180  
 agtaaaagaa tccaaacaaa atagagcaaa cactgtgccca tctgttttat accttactcc 240  
 agtaaatgca aactaataca caatgaaaaa aattacttat ttgagaactg gggagaggaa 300  
 ggagagggaa aggggtagat aaagaaaaga ggagagatta aaaggagcat aagaaaacct 360  
 cagagaataa taggtttgtg gtaaacatta ccgtggtaat gtttttaggg tatattcaca 420  
 tgtaaaaact tatccaatta tacattttta atatgtacag tttagtgtgt cagttatgcc 480  
 tctgtaaagt tgatttttaa aaaagtccta ttccaagtym acaatttcat ttg 533

<210> 43  
 <211> 480  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> allele  
 <222> 201  
 <223> 99-13929-201 : polymorphic base A or C

<220>  
 <221> misc\_binding

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$\langle 220 \rangle$

<221> primer\_bind  
<222> 109..127  
<223> 99-14021-108.mis complement

<220>  
<221> primer\_bind  
<222> 1..18  
<223> 99-14021.pu

<220>  
<221> primer\_bind  
<222> 460..477  
<223> 99-14021.rp complement

<400> 44  
tttgttggtta atcgcccctt ttctgcaaca cttgtgggtt agggaaaata attctaaagc 60  
aagagcaaag acagagttgg gagatcacca gtgaggttca attttcrtc acattcactc 120  
tgctccacac ctacagataat catgtgctta actgcgaaac ttgcttgaca attacagaac 180  
actttctcac ccattactac cttgatcctc acaattctgt ggggtagtag gagcagatgc 240  
tgaaattgcc atacgcaaat cagtgaactg aagcttagag acctccagca ggggcagagg 300  
gtcagcggaa actatcccag gggtcagcca acaagaaagt atattggaat cagagtatta 360  
aaataagaat aataaaacca actaaaattt accgtgcttt ttatttccac tcagtgccaa 420  
caattcttaa cagtgtcagt gatggatccc tgtgccccag gggacagact tcttact 477

<210> 45  
<211> 475  
<212> DNA  
<213> Homo sapiens

<220>  
<221> allele  
<222> 314  
<223> 99-14359-314 : polymorphic base G or C

<220>  
<221> misc\_binding  
<222> 291..337  
<223> 99-14359-314.probe

<220>  
<221> primer\_bind  
<222> 295..313  
<223> 99-14359-314.mis

<220>  
<221> primer\_bind  
<222> 315..333  
<223> 99-14359-314.mis complement

<220>  
<221> primer\_bind  
<222> 1..18  
<223> 99-14359.pu



<220>  
 <221> primer\_bind  
 <222> 457..475  
 <223> 99-14359.rp complement

<400> 45  
 ataaggggaat ggtgtgaggt gggaccagag gaggtgcac tgagaaagtg agaggggcaa 60  
 gacctcaggg gaagaaggga gggctgcacg gatgtctcag gcagagcagg cagcaccgga 120  
 aaaggtgggg gacactcctt ttggaccagc atataatttg gttaaagcct ctctgtttc 180  
 acctaataata taagcacatt tcaagataaa actactactt tattgtcatc aaatataaaa 240  
 gtaatttttt attcagggtt ttctaatact catctataaa ggcatttctt tcccacatgg 300  
 catgtgtttac aggstgttta acttaaagca attgtaaaag aaaagcctga agaaataagt 360  
 ctacaacgat ttacatcgtg tttatttttg tgtcaaaata tatgttaaaa tatacattag 420  
 ctatactaag ggaatcaaga gaagatcata attgctctta tgacttggga ttttag 475

<210> 46  
 <211> 473  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> allele  
 <222> 316  
 <223> 99-14364-415 : polymorphic base C or T

<220>  
 <221> misc\_binding  
 <222> 293..339  
 <223> 99-14364-415.probe

<220>  
 <221> primer\_bind  
 <222> 297..315  
 <223> 99-14364-415.mis

<220>  
 <221> primer\_bind  
 <222> 317..335  
 <223> 99-14364-415.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..19  
 <223> 99-14364.pu

<220>  
 <221> primer\_bind  
 <222> 453..473  
 <223> 99-14364.rp complement

<400> 46  
 gtgttttaat tcaaccagc tataagatac gaaatgatag aattgctcta gattctctat 60  
 tggttaaata aggagatatt tgtgctattg ccaataatac atgctgtacc tggataaacc 120  
 cctttgggca agttgtgatg caaataactca agaaaatag ccacatagtt acaacaggac 180



<212> DNA  
<213> Homo sapiens

<220>  
<221> allele  
<222> 412  
<223> 99-15229-412 : polymorphic base A or G

<220>  
<221> misc\_binding  
<222> 389..435  
<223> 99-15229-412.probe

<220>  
<221> primer\_bind  
<222> 393..411  
<223> 99-15229-412.mis

<220>  
<221> primer\_bind  
<222> 413..431  
<223> 99-15229-412.mis complement

<220>  
<221> primer\_bind  
<222> 1..20  
<223> 99-15229.pu

<220>  
<221> primer\_bind  
<222> 476..494  
<223> 99-15229.rp complement

<400> 48  
ctgtcattga gaaatgctac caataatact tagagaattt gatacaactc agtctgaaaa 60  
agctaagatt agcagaacag agctgtctcc aaatatattt agaactattt tatttaaggg 120  
attggaccca tttttgtatg tagttccaga ggagcagatg gtgaccactg tccaggcaga 180  
tgtgtctcaa tgtaaggaca acatctgtaa tattaataat tagaatgtat cctgtaattt 240  
tctctctacc cttggaaacc agtcgagatc cagagtcttt cactgggagg cttaaagcct 300  
agagcagcct tgggtgctaga ggcggacagg gataatgaac taatcttgaa ccaattcatc 360  
catagcaatc tcaatgcttt cgtttagctct tataggattt taatacggcc avaggaatga 420  
aggtagtctt gctggtttag aagccctgcc taccacaacc cctacaccac cccatcccct 480  
gcatagtctg atgt 494

<210> 49  
<211> 485  
<212> DNA  
<213> Homo sapiens

<220>  
<221> allele  
<222> 291  
<223> 99-15232-291 : polymorphic base G or T

<220>  
 <221> misc\_binding  
 <222> 268..314  
 <223> 99-15232-291.probe

<220>  
 <221> primer\_bind  
 <222> 272..290  
 <223> 99-15232-291.mis

<220>  
 <221> primer\_bind  
 <222> 292..310  
 <223> 99-15232-291.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..18  
 <223> 99-15232.pu

<220>  
 <221> primer\_bind  
 <222> 467..485  
 <223> 99-15232.rp complement

<400> 49  
 caatagaaca ggctgctcct ttataattat taatcatagt gtatattaat tcatcatcac 60  
 atacgtggct agaaaaaaaaa ttagaacaaa aagatatgtg atatgtaaag gcctacgata 120  
 attcagactt ctttgaggag agcttttatt ttattgttat tcttatttta tctcttgtca 180  
 atataaattg agagaataaa cagacaaaca ttacaaatta gtgattaatt gcatttaaag 240  
 cctagttaag actattttaag actatttatgc ataatacagg aaaactacct ktattattta 300  
 tagtgggtgc cttctgaagg atctgaagga gaatcagttc tatgcctctc tctcattcc 360  
 caggaggtgc ctggcattcc ttggcttgta gacgcacac cctaattctc acctctgcct 420  
 tcacatggtg tcccctgtgt gtgtgttttt gcccacatgtg tctcctcttt ttatatggat 480  
 gccag 485

<210> 50  
 <211> 464  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> allele  
 <222> 347  
 <223> 99-15241-347 : polymorphic base A or G

<220>  
 <221> misc\_binding  
 <222> 324..370  
 <223> 99-15241-347.probe

<220>  
 <221> primer\_bind  
 <222> 328..346

<223> 99-15241-347.mis

<220>

<221> primer\_bind

<222> 348..366

<223> 99-15241-347.mis complement

<220>

<221> primer\_bind

<222> 1..19

<223> 99-15241.pu

<220>

<221> primer\_bind

<222> 444..464

<223> 99-15241.rp complement

<400> 50

gttatgggtt	gaaaatctct	gagttcttgt	acatacaaaa	atctttactgt	tgtcacagtt	60
gaatcttagt	ttagatgggt	ataggatttt	tattcaaaaat	gcttttactc	cataagttta	120
aaaatattgt	tacattttcc	tcaagtatct	gatgttattg	atgagaagtt	taattctaata	180
ttgactcttg	ttcccttgta	ggtactatct	gttttccagt	ttgggaagct	tacatttctt	240
aaaattcaca	acataataatt	tacatactac	acaattcttt	ttaaagtata	caattcaatg	300
catttagtat	gttttagtac	atataactta	aattatgtat	atacaaratc	tctttataat	360
atgtgtagaa	tatgtagcat	attcacaaga	ttgttcaacc	atcaccactc	tctatttcca	420
gaatcttttc	ctccaaaaag	aaaccctgaa	cactatgatg	aata		464

<210> 51

<211> 550

<212> DNA

<213> Homo sapiens

<220>

<221> allele

<222> 196

<223> 99-15244-196 : polymorphic base A or G

<220>

<221> misc\_binding

<222> 173..219

<223> 99-15244-196.probe

<220>

<221> primer\_bind

<222> 177..195

<223> 99-15244-196.mis

<220>

<221> primer\_bind

<222> 197..215

<223> 99-15244-196.mis complement

<220>

<221> primer\_bind

<222> 1..20  
<223> 99-15244.pu

<220>  
<221> primer\_bind  
<222> 532..550  
<223> 99-15244.rp complement

<400> 51  
ctgcttctgg ttatgttttc ctaattgcc aatggtaaa aatgagaata atcattgaaa 60  
gagaaagcat aaagtagcaa aaatcctttc cagattaaaa aacgaagcaa agcatgtttc 120  
ccaagtaata atactctcat cttcctccct aatcctttac ccactacca gaagaagagt 180  
aaaatgtccg gatatrcttg aaggtaaaga tttctccttt taataaaatt agtcaccttg 240  
tacacatcag tagatcttga gaatgaaaag cttttctagt acattcattt caacctataa 300  
atgtttgact tttctctgtc attcatttac gacctgtgat cttttcattc cttttcagtt 360  
agaatatttt tcaaattttt attgatattt tctatttaac ccataggtta ttgggaaata 420  
cattgtttta tttctaatat atttgctttt ttttctactt atttcttttt ttcttaattc 480  
cacactggtc caaatatatt ctgcatatga tttaatat tttaatat taagttctgt agagactaac 540  
cttgtgccct 550

<210> 52  
<211> 452  
<212> DNA  
<213> Homo sapiens

<220>  
<221> allele  
<222> 404  
<223> 99-15252-404 : polymorphic base C or T

<220>  
<221> misc\_binding  
<222> 381..427  
<223> 99-15252-404.probe

<220>  
<221> primer\_bind  
<222> 385..403  
<223> 99-15252-404.mis

<220>  
<221> primer\_bind  
<222> 405..423  
<223> 99-15252-404.mis complement

<220>  
<221> primer\_bind  
<222> 1..18  
<223> 99-15252.pu

<220>  
<221> primer\_bind  
<222> 433..452  
<223> 99-15252.rp complement

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<400> 52
atgggggcat atagcaaccc tttagaaaca aaactacaaa aggtaagctt gtcttcttgc      60
atttcctttc tcttactaca tttaacatgg gaggttttct atgtctcaca ttcaaatatt    120
ctcactcggg ctgcctaatt ttccctgat ttcccatcac tctttatgaa ggcttgctac    180
tttagaatac acattttctt aacagaagat aataatcaga agatgtctcc caaatataag    240
tccaaatctt tcctatcatg ctgtgttctt tggctctttt gactttattt gaagtcagcc    300
ttgaagggga tagagatagg ctgtatgaag tccacgctga gaagttttgc cctgccctac    360
ttgtcctgta atatttcatg gatagcccag tggtgattaa accygtgtgt acaggaataa    420
ccatgagaat ttgttaaaaa tataggctct gg                                     452
```

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<210> 53
<211> 477
<212> DNA
<213> Homo sapiens
```

```
<220>
<221> allele
<222> 382
<223> 99-15253-382 : polymorphic base C or T
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```
<220>
<221> misc_binding
<222> 359..405
<223> 99-15253-382.probe
```

```
<220>
<221> primer_bind
<222> 363..381
<223> 99-15253-382.mis
```

```
<220>
<221> primer_bind
<222> 383..401
<223> 99-15253-382.mis complement
```

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<220>
<221> primer_bind
<222> 1..19
<223> 99-15253.pu
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<220>
<221> primer_bind
<222> 459..477
<223> 99-15253.rp complement
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<400> 53
aaaatcaatt ccccaacact cattttgtac gctaattttg taagatcctg aaaagtttca      60
ctattttatg gtttcatgtg ttacagatga aaaaaaaact agaattcaaa ttttctgagt    120
ttttttttac aatattttat gattacaaag ttagaagact aagaataaaa tggcctaatt    180
tccataatgt gagtggtaaa tgcagagcac tggcctaaag aaaatatttc aaaaaattag    240
tcactctttc cttaattttt ttccaacctg tgatctgttg aatgagcatt ttgcatatat    300
aaataaataa attactttgt aaataatctt gactggtttc tgttgaccac agtaaccac     360
tgcacagcac agcctgtaat tyctatgaac ctagggaat gtattttaagt ttattttttg    420
```

attacacagg tcctcattgt gtaactaaac attgcataga atatgccagt gatgatg 477

<210> 54  
 <211> 456  
 <212> DNA  
 <213> Homo sapiens  
 <220>  
 <221> allele  
 <222> 392  
 <223> 99-15256-392 : polymorphic base C or T

<220>  
 <221> misc\_binding  
 <222> 369..415  
 <223> 99-15256-392.probe

<220>  
 <221> primer\_bind  
 <222> 373..391  
 <223> 99-15256-392.mis

<220>  
 <221> primer\_bind  
 <222> 393..411  
 <223> 99-15256-392.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..18  
 <223> 99-15256.pu

<220>  
 <221> primer\_bind  
 <222> 439..456  
 <223> 99-15256.rp complement

<400> 54	
cctctctatg atgcttccta ttaagcaatt ggggaaatgt aataaacaag ggttggtgag	60
catcttcctt agtgagatgt ttttggaaga attggataat tgagtgaata atagtgagaa	120
actcctgtgt ctgatgttgc tccatgttgg aatgctttta tgttctcaga gaatgagtca	180
ctgagagcca attgtgatga tacacaatgg ttttaccag gttggatatg gtcctctgta	240
ctgggtaccct ttaagtcagt ggcactaatc agtcagtcac tgtcatgctt tgtgttggtc	300
catcatatgg tatgccctct tagagaacat cctgattagt ccttagacat cttttcaatt	360
tgaacactgg ggctcctcat tcgggtaaaa aytatggaca gtcagtgaac ctggttgcaat	420
ggcccctcat agcagattgg atctcaatgc actttg	456

<210> 55  
 <211> 501  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> allele



<222> 200  
<223> 99-15261-202 : polymorphic base A or G

<220>  
<221> misc\_binding  
<222> 177..223  
<223> 99-15261-202.probe

<220>  
<221> primer\_bind  
<222> 181..199  
<223> 99-15261-202.mis

<220>  
<221> primer\_bind  
<222> 201..219  
<223> 99-15261-202.mis complement

<220>  
<221> primer\_bind  
<222> 1..19  
<223> 99-15261.pu

<220>  
<221> primer\_bind  
<222> 481..501  
<223> 99-15261.rp complement

<400> 55  
cttctaattcc tttgtttcca cttattttat ttcattctctc attttatccc ttttttctaa 60  
attccattttt attatactta aggtgctttt aatatgggta tcatactcct gatagtgtta 120  
tttctttctt agtctttctta tataagcgct atacgttcac attccatctc ctttggttat 180  
ctttccattt cttcacccgar cctctttgct ctcttttttt atagctgggt cactcaaaat 240  
gtcttacttt gccatttttg aaatttatatt tcattctttt atgtactgaa taaaatttaa 300  
aaatacttta tcatgggtggg aggtacccgt gatgtccaaa taagtgttta tattaattgt 360  
tgggggtttt ttgtttgtgt gttttttgaa aggttaagaa aatctcattc agaaagtaag 420  
ttgtttaaaa attctggacc aaatttacca cacatcaagc agatacttac caagttgttt 480  
ggtagacatt agcagtattt a 501

<210> 56  
<211> 541  
<212> DNA  
<213> Homo sapiens

<220>  
<221> allele  
<222> 432  
<223> 99-15280-432 : polymorphic base C or T

<220>  
<221> misc\_binding  
<222> 409..455  
<223> 99-15280-432.probe

<220>  
 <221> primer\_bind  
 <222> 413..431  
 <223> 99-15280-432.mis

<220>  
 <221> primer\_bind  
 <222> 433..451  
 <223> 99-15280-432.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..18  
 <223> 99-15280.pu

<220>  
 <221> primer\_bind  
 <222> 521..541  
 <223> 99-15280.rp complement

<400> 56  
 atgtccatcc atcttgccca gagagagttt ctacaacact tcctctgcaa gccctttccc 60  
 tacttgccctc acctattgct ttcctctggt acgttgattt cccctcactg tttcttccaa 120  
 catcttccca cctcagagca tggacacttg ctgctctttc tgtgtcatga tgctgctcac 180  
 ttgtcccttt cttaatgtct cctccctgag ccaatcttct ccacccccac aacttacgca 240  
 cacttacatg tcatattttc cttcatagcc tttaacacca tttgaaatga tatatatttg 300  
 attgctttta aaatttctct gtccccccac taaatataaa cttcaggatg gcaagaatgt 360  
 agtccattat cttatttctc cagcctccat acttttaaga aaataaattt tggttgtata 420  
 agccatccag tyagtgttac ttgggtatag cacccttagc aaaagaatac aaaaaaaggg 480  
 agaatgtttg caatcatctg tttgaggcta ggaattccca gagagggaaa caaggagtaa 540  
 t 541

<210> 57  
 <211> 514  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> allele  
 <222> 428  
 <223> 99-15353-428 : polymorphic base C or T

<220>  
 <221> misc\_binding  
 <222> 405..451  
 <223> 99-15353-428.probe

<220>  
 <221> primer\_bind  
 <222> 409..427  
 <223> 99-15353-428.mis

<220>  
 <221> primer\_bind

<222> 429..447  
<223> 99-15353-428.mis complement

<220>  
<221> primer\_bind  
<222> 1..18  
<223> 99-15353.pu

<220>  
<221> primer\_bind  
<222> 495..514  
<223> 99-15353.rp complement

<400> 57  
tgggaatgga ggtagtagac gatgaggtct ccaccctctg actttgcaga gatgggcaag 60  
gccaagtgtt ggaagggctt aaacacacac cggagtattc tgtgagaacc agtggatttc 120  
agaggatggc aatgacacca cttgccttct gcctcaggag gataactgat ggccgtgtgt 180  
gggatgcact ggagagcaag agctggcttg cagggagacc agctggatga ttttctttca 240  
tttattttat tcattcaaca cacattcatc tggggttcac tctgtgcccc acactgggca 300  
tttccaaata gtccagatgg cagtaagcat ggttgtggca gtaggaatgg gaaggctggg 360  
aggggtatga gaggcattac aaacgggaag tgggagtggc accccagaaa agtctagttt 420  
aaggtgcyag tggatgtgtg catgtgtgcg cgggggtgtc tagagggtgg cgggcagctg 480  
gaaattgagg tcaagtgcct aaagaacaac tcgt 514

<210> 58  
<211> 489  
<212> DNA  
<213> Homo sapiens

<220>  
<221> allele  
<222> 150  
<223> 99-15355-150 : polymorphic base C or T

<220>  
<221> misc\_binding  
<222> 127..173  
<223> 99-15355-150.probe

<220>  
<221> primer\_bind  
<222> 131..149  
<223> 99-15355-150.mis

<220>  
<221> primer\_bind  
<222> 151..169  
<223> 99-15355-150.mis complement

<220>  
<221> primer\_bind  
<222> 1..18  
<223> 99-15355.pu

<220>  
 <221> primer\_bind  
 <222> 471..489  
 <223> 99-15355.rp complement

<400> 58  
 taactttctcc gtctctcctt cttagcccat atgtcaataa tgactgaaag tattcatttc 60  
 catcttttaa ctgcctattc cagccacctc ccacctccat ctctttcctt ctaagttttc 120  
 ttcattcttct actttgggca aaaggaaaty gatgtgtcag acaggcctag ttttgaattc 180  
 tggatctgct agcacttctc tgtgtgtcct tggttatatg atatagtctt aaaccttaat 240  
 gttcttgcct gtaaaatggg gataataaaa acctcttaac agtgggtgtt tcatgcagct 300  
 ttcattacaa acttcctcat tcaaaatctt caatgatttc catttttcac aaaatgaaat 360  
 tcaaaatttc tgtagattat tgagacaagt cccctactct tcacctaaat ttatctttta 420  
 tttattctct catcattatc aacaactact aggctttgtt gccttgactc cagaggcaaa 480  
 aatcttattc 489

<210> 59  
 <211> 468  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> allele  
 <222> 227  
 <223> 99-15685-227 : polymorphic base A or G

<220>  
 <221> misc\_binding  
 <222> 204..250  
 <223> 99-15685-227.probe

<220>  
 <221> primer\_bind  
 <222> 208..226  
 <223> 99-15685-227.mis

<220>  
 <221> primer\_bind  
 <222> 228..246  
 <223> 99-15685-227.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..18  
 <223> 99-15685.pu

<220>  
 <221> primer\_bind  
 <222> 449..468  
 <223> 99-15685.rp complement

<400> 59  
 aaacaaaggc acgcagagga taaggcatga gtccaaccag cagcatctcc ctcccgaatg 60  
 agtacagaaa tgatcaatac tcgaagagaa aaagatgctt tcagtgtgct ttacctgaaa 120

acttccttaa	gcagcttcac	tttattgtca	ggatatcgct	ttgtgtttgt	atcatctaag	180
aaagctcgcg	catatgctag	tgggccagca	ttgacctaga	caaagarcaa	agattttcag	240
ttccactagg	aagaaaatca	ccatgaccat	ctgctcagtt	tcagtttgca	ggcactaaaa	300
agcccgttcg	cgtgagctac	tcacaatccc	tgccttccag	gaacttaagc	ccaaaaagaa	360
accacaaagc	tcactctgtt	gcacaccact	tgattccatg	atctcagcca	tcttcagggc	420
acttgtgatg	atggtttact	ttatgtaaga	agaaaccaat	gcttgga		468

<210> 60  
 <211> 500  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> allele  
 <222> 428  
 <223> 99-15695-428 : polymorphic base C or T

<220>  
 <221> misc\_binding  
 <222> 405..451  
 <223> 99-15695-428.probe

<220>  
 <221> primer\_bind  
 <222> 409..427  
 <223> 99-15695-428.mis

<220>  
 <221> primer\_bind  
 <222> 429..447  
 <223> 99-15695-428.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..18  
 <223> 99-15695.pu

<220>  
 <221> primer\_bind  
 <222> 481..500  
 <223> 99-15695.rp complement

<400> 60	
atcagccttt	gtgaggagga ggcctgcct gctctcctcc tgagctgatg ggtcagtcac 60
accaggacaa	aggtctgccc ggggctgtgt gggttcctcc ttcttgagct gcacaccagc 120
atctgctgaa	caccttctgg agctcagctc agtgtctcgt ccagagacac tggttccctt 180
ggcttctcag	caactctcgg atctgggcct ggggtctaacc tcagcgggtgg tcttgcccat 240
ttctagggcc	tcacaattca gcctcatgtc ttcacctgtg gctcttttgc aaggctcaga 300
aagctctagg	gtcagttcca gatgactccc accagcatgc cagtaggagc caccaccccc 360
tctcagccag	cgccaccata ttccaggcaa attccaactg acacagactt caaggaacga 420
ttgtagcygt	tgttcttctg tcttccaaat ggaagagtgc attattgggg tcccttctag 480
cacgcatttc	attccccacc
	500

<210> 61

<211> 472  
<212> DNA  
<213> Homo sapiens

<220>  
<221> allele  
<222> 310  
<223> 99-15703-310 : polymorphic base C or T

<220>  
<221> misc\_binding  
<222> 287..333  
<223> 99-15703-310.probe

<220>  
<221> primer\_bind  
<222> 291..309  
<223> 99-15703-310.mis

<220>  
<221> primer\_bind  
<222> 311..329  
<223> 99-15703-310.mis complement

<220>  
<221> primer\_bind  
<222> 1..18  
<223> 99-15703.pu

<220>  
<221> primer\_bind  
<222> 452..472  
<223> 99-15703.rp complement

<400> 61  
agggtcttgg gttataggcg ctgaatttct tctaaagcta acctgactct gatgctagaa 60  
gagcccattt aaggaaagaa aaacactttt cattgctcga tcaaagttca tccatttttg 120  
aaaagacatc aaaccaagtg tgtgacacca ggcaccata tccttcctct tccccaccac 180  
cccacccctg tcctcagggc agtgacagtg aagcctgggtg caggtcccgc tgctgctttt 240  
tgaagtggca catgctttat tttcttaaaa agaagtgaga gacaacctat gctacaggag 300  
gctctgtgay gtttttctga agtacaaccc cttgctctgc cagggcagct gtaaagggtc 360  
taaagagccc tgagaaagga gagaggattt gggaagccga ggaggcagag ggagaccaca 420  
tagcacatgg agttctgaaa gggcccaagt ggagacagaa aacgagtcac gt 472

<210> 62  
<211> 470  
<212> DNA  
<213> Homo sapiens

<220>  
<221> allele  
<222> 400  
<223> 99-15870-400 : polymorphic base A or G

<220>  
 <221> misc\_binding  
 <222> 377..423  
 <223> 99-15870-400.probe

<220>  
 <221> primer\_bind  
 <222> 381..399  
 <223> 99-15870-400.mis

<220>  
 <221> primer\_bind  
 <222> 401..419  
 <223> 99-15870-400.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..21  
 <223> 99-15870.pu

<220>  
 <221> primer\_bind  
 <222> 452..470  
 <223> 99-15870.rp complement

<400> 62  
 gctcaaatgt atcaaacaca gtttctgtgg tcaagttcct ctccctttct aaatttgctt 60  
 agaggatctc ataaaacgta actcctctga caagggaacc atttttagcac caacactgca 120  
 aaagcttctg tgttcctaag ggaaagatcc tttcctgaat taaatttaac ctcttttagta 180  
 ctcccattta gccacctgat aaatccactt gagctatctt ttgggaagag agaggatatct 240  
 gggaacaata acacttcctt tttgaacagt ttaataaagc tttgtgagat ttcaagatga 300  
 aagataatgt gtaatgctga tagtgccctc caaggctctg cattcatgga tccaattacg 360  
 ttttttgtca tggtaaaagc cacagtggat atattaaatr agagtgtggt ttaagaatga 420  
 aggcccagga gtctggagat ctggtttcta aggctgactt cacttctgct 470

<210> 63  
 <211> 469  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> allele  
 <222> 287  
 <223> 99-16321-287 : polymorphic base A or C

<220>  
 <221> misc\_binding  
 <222> 264..310  
 <223> 99-16321-287.probe

<220>  
 <221> primer\_bind  
 <222> 268..286  
 <223> 99-16321-287.mis

<220>  
 <221> primer\_bind  
 <222> 288..306  
 <223> 99-16321-287.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..20  
 <223> 99-16321.pu

<220>  
 <221> primer\_bind  
 <222> 451..469  
 <223> 99-16321.rp complement

<400> 63  
 ctttaggaat atcccttctg atttgaacaa cattttgcta tccaagttct gtctactttt 60  
 ttaacaagtt cttgctccgt gtgtctcctt ttgcttggtc tcaagtaagg gagtaacagg 120  
 gataaactcc cactccttgg taaatctttc tatcattttt ggaaatctca tccattgtag 180  
 taaatgctct taaatcttca tcttcaggcc gtgacttcca tctagcctcc attcacgttt 240  
 ccgggtttat gtctgcaatg agcattccgt ggctctacat agatgcmcca ccatacctag 300  
 aacccatgta tcccaaaactc aattctttct ttcccaggac attacttctt gcacttcctt 360  
 agtctatcaa tggcactggt attctcttga ccatctagac ttgaaatttt ggggtttgga 420  
 ctctctctgc tcccttgctt tatatgtaat cagacatcaa gtctcaatc 469

<210> 64  
 <211> 544  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> allele  
 <222> 194  
 <223> 99-16333-194 : polymorphic base A or G

<220>  
 <221> misc\_binding  
 <222> 171..217  
 <223> 99-16333-194.probe

<220>  
 <221> primer\_bind  
 <222> 175..193  
 <223> 99-16333-194.mis

<220>  
 <221> primer\_bind  
 <222> 195..213  
 <223> 99-16333-194.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..19



66210T"48E9T460

<223> 99-16333.pu

<220>

<221> primer\_bind

<222> 524..544

<223> 99-16333.rp complement

<400> 64

atttacc	ccg	tctgc	cttgc	aatttc	cagga	tcagt	tataca	tcaaat	caag	tgaaca	accc	60
aggga	attct	gccgt	tacct	tttaga	aaaca	gaataa	atat	taacag	agct	ttactt	cttt	120
ccacca	agga	ggact	atatg	ttaata	acagt	aattta	caact	ggaaaa	aaata	taaatg	aaag	180
ggttt	tagaac	ctcrta	aactt	taaaaa	ataac	ataatt	cctc	ctaga	acatt	cctttc	actt	240
gtgatt	ctca	aagcact	ttg	catttcc	cag	ctattg	gcag	ggctg	gaatt	aggatc	aaag	300
tatcact	aaaa	tggtag	gtga	aataaat	gtg	aagctg	at	tcagg	agtac	aggaat	ggag	360
tcatc	aggcg	actttta	aaagt	taagaat	ctg	ttggag	cagc	tgcca	ataaa	tcaagg	ccca	420
aaggag	aaaag	ttcttt	ggaa	accttg	aaat	attgtat	acata	tttag	ataat	tattgt	tgtt	480
gtcaat	gtta	acgaaaa	aaag	caataa	atca	gggag	atggc	actgat	gagt	gaggag	aaat	540
agac												544

<210> 65

<211> 475

<212> DNA

<213> Homo sapiens

<220>

<221> allele

<222> 149

<223> 99-5873-159 : polymorphic base C or T

<220>

<221> misc\_binding

<222> 126..172

<223> 99-5873-159.probe

<220>

<221> primer\_bind

<222> 130..148

<223> 99-5873-159.mis

<220>

<221> primer\_bind

<222> 150..168

<223> 99-5873-159.mis complement

<220>

<221> primer\_bind

<222> 1..18

<223> 99-5873.pu

<220>

<221> primer\_bind

<222> 457..475

<223> 99-5873.rp complement

552707"43E9T460

<220>  
<221> misc\_feature  
<222> 409  
<223> n=a, g, c or t

<400> 65  
gcgtaacaat aagcaggggt agtcgccaca aaacttgaga taagaggaaa actaaaaaag 60  
tctaataaaa tcagtagtct taaaaagatg acatgatagg aagagaagtg ttaaaaaaga 120  
aaaaaaatag gtatgaaaga gagtaacaya taccggaaaa gggataaaat acatcctttg 180  
aaagaacaaa gagttattca aattgaattc ttaatgaatt acttaaacag cagattagat 240  
attgttaaaa agaggaatag ggaattaaat gatatatgtg atgatattac ctagtgtaac 300  
catcaaagat gtattgcaaa tgataaagaa aaaaatgctg ccatggcaat attaatatca 360  
taaaaatata ctttaagaag taaataaatg caactaggaa tagagaaans dvhatgaata 420  
ataatattta amaaavvgta taacaagtat acataagatg taatatccta aaccg 475

<210> 66  
<211> 511  
<212> DNA  
<213> Homo sapiens

<220>  
<221> allele  
<222> 49  
<223> 99-5912-49 : polymorphic base A or G

<220>  
<221> misc\_binding  
<222> 26..72  
<223> 99-5912-49.probe

<220>  
<221> primer\_bind  
<222> 30..48  
<223> 99-5912-49.mis

<220>  
<221> primer\_bind  
<222> 50..68  
<223> 99-5912-49.mis complement

<220>  
<221> primer\_bind  
<222> 11..31  
<223> 99-5912.pu

<220>  
<221> primer\_bind  
<222> 494..511  
<223> 99-5912.rp complement

<400> 66  
aaatataata gtcaaatcat gttaccatta ggacacatta aaaatgtcra attaccttgg 60  
gaccttatat gaacatatta agataataat gatagtgttc agtgcaatat tcagatcaat 120  
agtttaaacc caaaatattt ataccttcag attagatgta tgcaaagtca ttgattcatg 180

tgtcttttat	ctgttggtta	catttgagga	aatatttgag	aaatatttca	aaatggaatt	240
tatataaatt	taaacacata	atgggtttat	gtaaaaatat	tgctaaatta	cattttcccc	300
ttaattctta	tttcttgga	acgtgcctta	gtcgtgaaa	tattcataca	ttaacacaat	360
gaaagaagt	aaccttacta	ggctttgact	atcagggttg	ctgttggttt	ttgactattg	420
tgaaactata	gcctgatttc	taaatcagga	agaaacgtgt	attgttggtta	atatggacac	480
atgacatatt	tgtctgcctg	acttttgatc	c			511

<210> 67  
 <211> 485  
 <212> DNA  
 <213> Homo sapiens

<220>  
 <221> allele  
 <222> 210  
 <223> 99-6012-220 : polymorphic base G or T

<220>  
 <221> misc\_binding  
 <222> 187..233  
 <223> 99-6012-220.probe

<220>  
 <221> primer\_bind  
 <222> 191..209  
 <223> 99-6012-220.mis

<220>  
 <221> primer\_bind  
 <222> 211..229  
 <223> 99-6012-220.mis complement

<220>  
 <221> primer\_bind  
 <222> 1..19  
 <223> 99-6012.pu

<220>  
 <221> primer\_bind  
 <222> 467..485  
 <223> 99-6012.rp complement

<400> 67	
gtcttgactt	gttttcctga
gtcttctcag	ttttaataat
ctgctcccag	gtcagtcctg
atttcctgat	tttagagatt
gcataaaaaa	atgtacagca
caacaaagac	cacatggagt
tacttagtgt	tttatttcca
tttagttgtt	tctaaagtag
ggata	
gggtccaggt	tgatttgcac
ccctgtgggt	tctcaggacc
ggttcccttg	tggtctcatc
aattttgaag	actttccttc
atggaatgag	atatactctg
agtttcctcg	aaaggcacca
atgtgaagta	tttggctgtc
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gtttattttt aatttcaaaa atgtactgca taaatgcagc aaaaccttta ttgtcacata 360  
ttaaacaatg tacattattg tgtgcaaatt aaaatttcat taccttaaac caaaaagtga 420  
gttggccaga tagtaaataa tttaggctct aaggctgaaa agcgcttgta ttaattactc 480  
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<212> DNA  
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 <223> 99-7308-157.mis

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 ggtatatgta ttcattggtga caaatatttatt cacatyttaa gtaattttta gtaattcaca 180  
 ttttaagtaa ttttcctgaa tgtgcctcat tggcttctgt gcctcttcag aaaagatgaa 240  
 ctaaactctg gcatatgtgt tcagatttca acattccggt gttttcattg tggataattt 300  
 ctgtcccata tttttgtgta aagtttagaca ataaagtgtt aatattctgg cgtcggcaca 360  
 ttttctttcc tgataaataa caattcacat atctttttta aatatcagag aatatagtaa 420  
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 ggcaactat 489

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<223> sequencing oligonucleotide PrimerRP

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18

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